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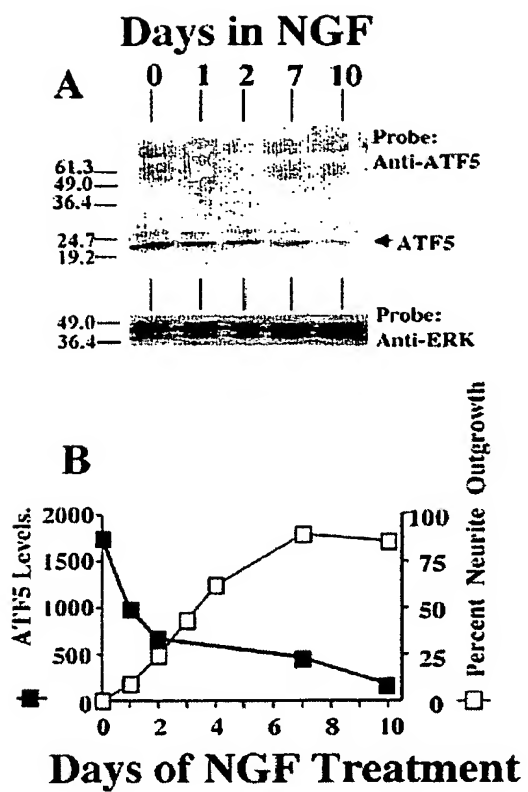


Fig. 1

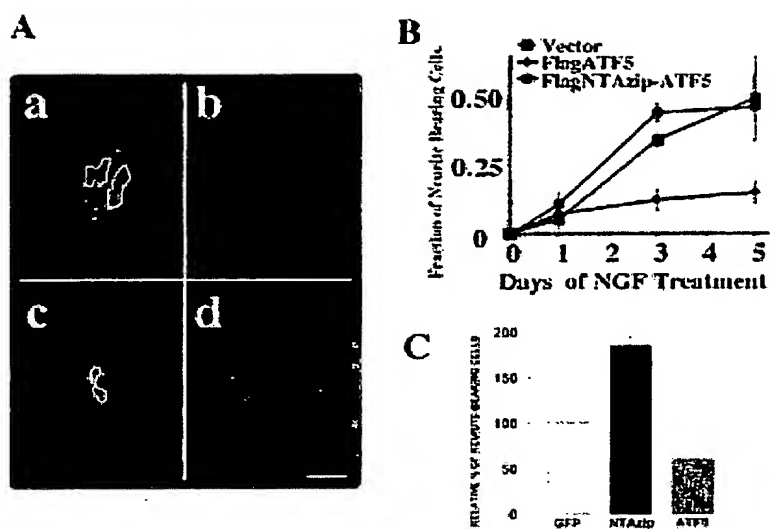


Fig. 2

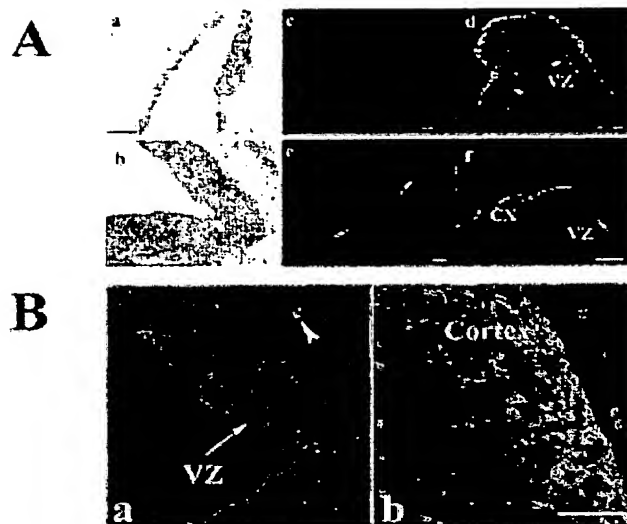


Fig. 3

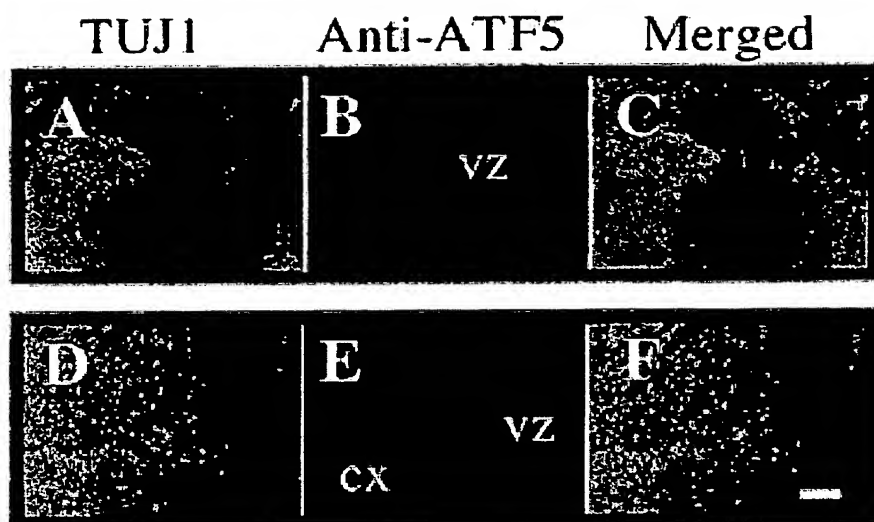


Fig. 4

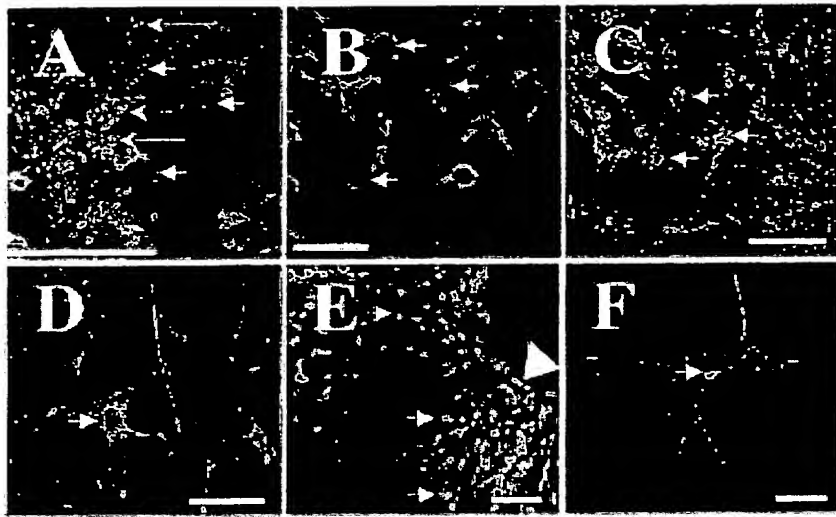


Fig. 5

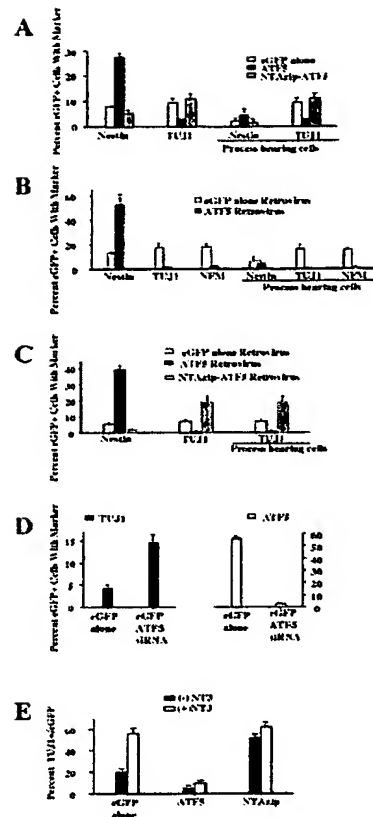


Fig. 6

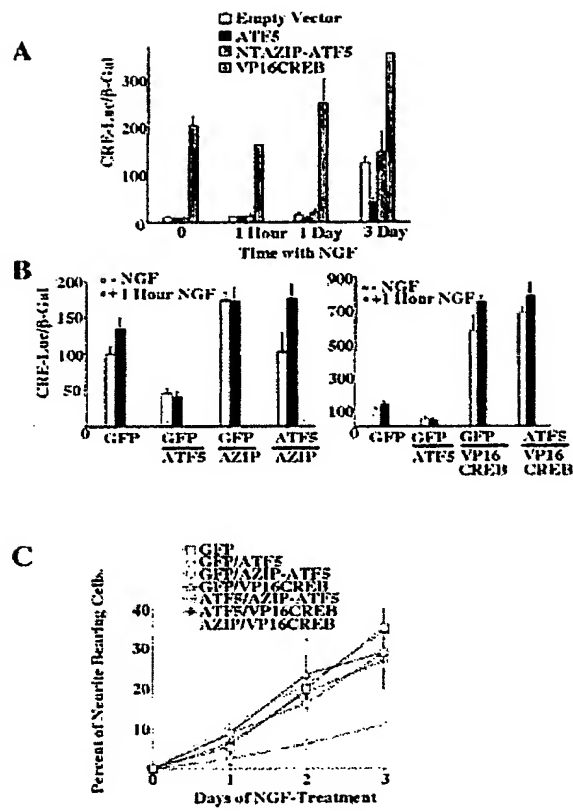


Fig. 7

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 121 cccccctggg cccctatgag gtcctgggg gtgccctgga gggcgggctt ccaggggggg
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 661 accagaataa gtcagctgct ctcaggtacc gccagaggaa gcgggcagag ggcgagggcc
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 1021 agaaccaagc tcga

Fig. 8

MSLLATLGLELDRALLPASGLGWLVDYGKLPLAPAPLGPYEVLGGALEGGLP
GGGEPLAGDGFSDWMTERVDFTALLPLEAPLPPGTLPPPSPAPPDLEAMAS
LLKKELEQMEDFFLDAPLLPPPSPPPPPPPPAPSLPLPLPLPTFDLPQPPTLDTL
DLLAVYCRSEAGPGDSGLTTLVPVQQPPPLAPLPSPSRPAPYPSPASTRGDR
KQKKRDQNKSAALRYRQRKRAEGEAGEGECQGLEARNRELRERAESVEREI
QYVKDLLIEVYKARSQRTRSA

Fig. 9